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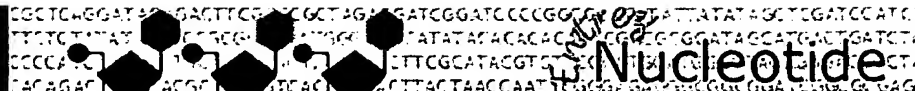
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Search for [Limits](#)[Preview/Index](#)[History](#)[Clipboard](#)[Details](#)Display Show Range: from to ☐ Reverse complemented strand Features: ☐ SNP ☐ CDD ☐1: [AF081469](#). Reports *Stachybotrys char...*[gi:3420911][Links](#)

- [Features](#)
- [Sequence](#)

LOCUS AF081469 936 bp DNA linear PLN 11-JUN-1999

DEFINITION *Stachybotrys chartarum* strain UAMH 7900 18S ribosomal RNA gene, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA gene and internal transcribed spacer 2, complete sequence; and 28S ribosomal RNA gene, partial sequence.

ACCESSION AF081469

VERSION AF081469.1 GI:3420911

KEYWORDS .

SOURCE *Stachybotrys chartarum*

ORGANISM *Stachybotrys chartarum*

Eukaryota; Fungi; Ascomycota; mitosporic Ascomycota; *Stachybotrys*.

REFERENCE 1 (bases 1 to 936)

AUTHORS Haugland, R.A. and Heckman, J.L.

TITLE Identification of putative sequence specific PCR primers for detection of the toxigenic fungal species *Stachybotrys chartarum*

JOURNAL Mol. Cell. Probes 12 (6), 387-396 (1998)

PUBMED 9843656

REFERENCE 2 (bases 1 to 936)

AUTHORS Haugland, R.A. and Heckman, J.L.

TITLE Direct Submission

JOURNAL Submitted (30-JUL-1998) Microbiological & Chemical Exposure Assessment Research Division, National Exposure Research Laboratory, U.S. Environmental Protection Agency, 26 W. Martin Luther King Dr., Cincinnati, OH 45268, USA

FEATURES Location/Qualifiers

source 1..936

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/db_xref="taxon:74722"

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misc RNA 362..520

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/product="5.8S ribosomal RNA"

misc RNA 678..855

/product="internal transcribed spacer 2"

/note="ITS2"

rRNA 856..>936

/product="28S ribosomal RNA"

ORIGIN

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